

SEQUENCE LISTING

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<120> CRYSTAL STRUCTURE OF BACE AND USES THEREOF

<130> 16163-015001

<140> US 09/955,737

<141> 2001-09-19

<150> US 60/234,576

<151> 2000-09-22

<160> 15

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 501

<212> PRT

<213> Homo sapiens

180

<400> 1

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Glu Gly Ile Leu Gly Leu Ala Tyr Ala Glu Ile Ala Arg Pro Asp Asp

185

170

190

Ser Leu Glu Pro Phe Phe Asp Ser Leu Val Lys Gln Thr His Val Pro

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200
Asn Leu Phe Ser Leu Gln Leu Cys Gly Ala Gly Phe Pro Leu Asn Gln
Ser Glu Val Leu Ala Ser Val Gly Gly Ser Met Ile Ile Gly Gly Ile
                    230
                                         235
Asp His Ser Leu Tyr Thr Gly Ser Leu Trp Tyr Thr Pro Ile Arg Arg
                                     250
Glu Trp Tyr Tyr Glu Val Ile Ile Val Arg Val Glu Ile Asn Gly Gln
                                 265
Asp Leu Lys Met Asp Cys Lys Glu Tyr Asn Tyr Asp Lys Ser Ile Val
                            280
                                                 285
Asp Ser Gly Thr Thr Asn Leu Arg Leu Pro Lys Lys Val Phe Glu Ala
                        295
Ala Val Lys Ser Ile Lys Ala Ala Ser Ser Thr Glu Lys Phe Pro Asp
                    310
                                         315
Gly Phe Trp Leu Gly Glu Gln Leu Val Cys Trp Gln Ala Gly Thr Thr
                325
                                     330
Pro Trp Asn Ile Phe Pro Val Ile Ser Leu Tyr Leu Met Gly Glu Val
                                 345
Thr Asn Gln Ser Phe Arg Ile Thr Ile Leu Pro Gln Gln Tyr Leu Arg
                             360
                                                 365
Pro Val Glu Asp Val Ala Thr Ser Gln Asp Asp Cys Tyr Lys Phe Ala
                        375
                                             380
Ile Ser Gln Ser Ser Thr Gly Thr Val Met Gly Ala Val Ile Met Glu
                    390
                                         395
Gly Phe Tyr Val Val Phe Asp Arg Ala Arg Lys Arg Ile Gly Phe Ala
                                     410
Val Ser Ala Cys His Val His Asp Glu Phe Arg Thr Ala Ala Val Glu
            420
                                 425
                                                     430
Gly Pro Phe Val Thr Leu Asp Met Glu Asp Cys Gly Tyr Asn Ile Pro
                            440
Gln Thr Asp Glu Ser Thr Leu Met Thr Ile Ala Tyr Val Met Ala Ala
                        455
                                             460
Ile Cys Ala Leu Phe Met Leu Pro Leu Cys Leu Met Val Cys Gln Trp
                    470
                                        475
Arg Cys Leu Arg Cys Leu Arg Gln Gln His Asp Asp Phe Ala Asp Asp
                485
                                    490
Ile Ser Leu Leu Lys
            500
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<213> Homo sapiens
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Ala Leu Glu Val Pro Thr Asp Gly Asn Ala Gly Leu Leu Ala Glu Pro
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Gln Ile Ala Met Phe Cys Gly Arg Leu Asn Met His Met Asn Val Gln
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Asn Gly Lys Trp Asp Ser Asp Pro Ser Gly Thr Lys Thr Cys Ile Asp

Thr Lys Glu Gly Ile Leu Gln Tyr Cys Gln Glu Val Tyr Pro Glu Leu

Gln	Ile	Thr	Asn	Val 85	Val	Glu	Ala	Asn	Gln 90	Pro	Val	Thr	Ile	Gln 95	Asn
Trp	Сув	Lys			Arg	Lys	Gln	Cys 105		Thr	His	Pro	His 110		Val
Ile	Pro	_	100 Arg	Cys	Leu	Val	Gly		Phe	Val	Ser	_		Leu	Leu
Val	Pro	115 Asp	Lys	Cys	Lys	Phe	120 Leu	His	Gln	Glu	Arg	125 Met	Asp	Val	Cys
	130					135					140				
Glu	Thr	His	Leu	His	Trp	His	Thr	Val	Ala	Lys	Glu	Thr	Cys	Ser	Glu
145					150					155			-		160
	Ser	Thr	Asn	Leu 165		Asp	Tyr	Gly	Met 170		Leu	Pro	Cys	Gly 175	
Asp	Lys	Phe	_		Val	Glu	Phe			Cys	Pro	Leu			Glu
_	_	_	180	_	_		_	185			_	_	190	_	
Ser	Asp	Asn 195	Val	Asp	Ser	Ala	Asp 200	Ala	Glu	Glu	Asp	Asp 205	Ser	Asp	Val
Trp	Trp	Gly	Gly	Ala	Asp	Thr	Asp	Tyr	Ala	Asp	Gly	Ser	Glu	Asp	Lys
	210					215					220				
Val	Val	Glu	Val	Ala	Glu	Glu	Glu	Glu	Val	Ala	Glu	Val	Glu	Glu	Glu
225					230					235					240
	. דת	7.00	7 an	7 an		7 00	Asp	C1	7 an		7 an	C1.,	1701	C1	
Gru	AIA	Asp	Asp		Gru	Asp	Asp	GIU		СТУ	Asp	GIU	vai		GIU
~-3		~-7	~-7	245	_		~-3		250	~-7	_	_,	_,	255	
GIu	Ala	GIu		Pro	Tyr	Glu	Glu		Thr	GIU	Arg	Thr		ser	пте
			260					265					270		
Ala	Thr	Thr	Thr	Thr	Thr	Thr	Thr	Glu	Ser	Val	Glu	Glu	Val	Val	Arg
		275					280					285			
Val	Pro	Thr	Thr	Ala	Ala	Ser	Thr	Pro	Asp	Ala	Val	Asp	Lys	Tyr	Leu
	290					295			_		300	_	-	-	
Glu	Thr	Pro	Glv	Asp	Glu	Asn	Glu	His	Ala	His	Phe	Gln	Lvs	Ala	Lvs
.305			1		310					315			-7-		320
	λκα	T.OU	Glu	715		uic	Arg	Glu	720		cor	Gln	Val	Mot	
Giu	ALG	пец	Giu	325	цуь	птъ	Arg	Giu	330	MEC	SCI	GIII	vai	335	Arg
a 1		~ 1	~ 1		~ 1	3	G 3			.	T	D	.		
GIU	Trp	GIU		Ата	GIU	Arg	Gln		гуѕ	ASI	ьeu	Pro	-	Ата	Asp
	_		340					345		_			350		
Lys	Lys		Val	Ile	Gln	His	Phe	Gln	Glu	Lys	Val		Ser	Leu	Glu
		355					360					365			
Gln	Glu 370	Ala	Ala	Asn	Glu	Arg 375	Gln	Gln	Leu	Val	Glu 380	Thr	His	Met	Ala
Arq	Val	Glu	Ala	Met	Leu	Asn	Asp	Arq	Arq	Arq	Leu	Ala	Leu	Glu	Asn
385					390		_	_		395					400
Tvr	Ile	Thr	Ala	Leu	Gln	Ala	Val	Pro	Pro	Ara	Pro	Ara	His	Val	Phe
4				405					410	3		5		415	
Δan	Mot	T.011	Lare		Tir	T = V	Arg	λΊэ		Gln	Lare	Nen	λκα		Uic
Pon	MCC	БСи		цуз	TYL	vai	Arg		Giu	GIII	цуз	Asp		GIII	птэ
m1	.	T	420	-1	~ 7	'		425			_	_	430	_	~ 7
Thr	ьeu		HIS	Pne	GIu	His	Val	Arg	Met	vaı	Asp		ьуs	ьуs	Ala
		435					440					445			
Ala	Gln	Ile	Arg	Ser	Gln	Val	Met	Thr	His	Leu	Arg	Val	Ile	Tyr	Glu
	450					455					460				
Arg	Met	Asn	Gln	Ser	Leu	Ser	Leu	Leu	Tyr	Asn	Val	Pro	Ala	Val	Ala
465					470				-	475					480
	Glu	Ile	Gln	Asp	Glu	Val	Asp	Glu	Leu	Leu	Gln	Lvs	Glu	Gln	
				485		- ~ -			490	u		-10	J_ u	495	
Ф~	C-~	7 c.~	λ α∽		T 033	ת דת	7.~~	Mo÷		C~~	C1	Dro	7~~		0
тйт	261	wsh		vaı	ьeu	HIG	Asn		TIG	261	GIU	PIO	_	TTE	Set.
m	03 .	7. –	500	7 T	. -	N/ - ·	D	505	.	m7.	α·1		510	m¹-	m¹-
Tyr	GTA		Asp	АТА	ьeu	met	Pro	ser	ьeu	Inr	GIU		гÀг	Thr	Thr
		515	_	_			520			_		525			
Val	Glu	Leu	Leu	Pro	Val	Asn	Gly	Glu	Phe	Ser	Leu	Asp	Asp	Leu	Gln

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530
                        535
                                            540
Pro Trp His Ser Phe Gly Ala Asp Ser Val Pro Ala Asn Thr Glu Asn
                    550
                                        555
Glu Val Glu Pro Val Asp Ala Arg Pro Ala Ala Asp Arg Gly Leu Thr
                                     570
Thr Arg Pro Gly Ser Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile Ser
                                585
            580
Glu Val Lys Met Asp Ala Glu Phe Arg His Asp Ser Gly Tyr Glu Val
                            600
His His Gln Lys Leu Val Phe Phe Ala Glu Asp Val Gly Ser Asn Lys
                        615
                                            620
Gly Ala Ile Ile Gly Leu Met Val Gly Gly Val Val Ile Ala Thr Val
                    630
                                        635
Ile Val Ile Thr Leu Val Met Leu Lys Lys Gln Tyr Thr Ser Ile
                                     650
His His Gly Val Val Glu Val Asp Ala Ala Val Thr Pro Glu Glu Arg
His Leu Ser Lys Met Gln Gln Asn Gly Tyr Glu Asn Pro Thr Tyr Lys
                            680
Phe Phe Glu Gln Met Gln Asn
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<213> Artificial Sequence
<220>
<223> APP inhibitor peptide
<220>
<221> VARIANT
<222> 5
<223> Xaa = Sta = Statine
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Ser Glu Val Asn Xaa Val Ala Glu Phe
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<211> 29
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<213> Artificial Sequence
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<223> Primer
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gctctagaac ccagcacggc atccggctg
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<211> 42
<212> DNA
<213> Artificial Sequence
<220>
<223> Primer
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42

<400> 5

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<223> Synthetically generated peptide
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Ser Glu Val Asn Leu Asp Ala Glu Phe Arg
<210> 7
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Gly Tyr Tyr Val Glu Met Thr Val Gly Ser Pro Pro Gln Thr Leu Asn
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                                25
Ile Leu Val Asp Thr Gly Ser Ser Asn Phe Ala Val Gly Ala Ala Pro
His Pro Phe Leu His Arg Tyr Tyr Gln Arg Gln Leu Ser Ser Thr Tyr
Arg Asp Leu Arg Lys Gly Val Tyr Val Pro Tyr Thr Gln Gly Lys Trp
                    70
                                         75
Glu Gly Glu Leu Gly Thr Asp Leu Val Ser Ile Pro His Gly Pro Asn
                                    90
Val Thr Val Arg Ala Asn Ile Ala Ala Ile Thr Glu Ser Asp Lys Phe
                                105
Phe Ile Asn Gly Ser Asn Trp Glu Gly Ile Leu Gly Leu Ala Tyr Ala
                            120
                                                 125
Glu Ile Ala Arg Pro Asp Asp Ser Leu Glu Pro Phe Phe Asp Ser Leu
                        135
Val Lys Gln Thr His Val Pro Asn Leu Phe Ser Leu Gln Leu Cys Gly
                    150
                                         155
Ala Gly Phe Pro Leu Asn Gln Ser Glu Val Leu Ala Ser Val Gly Gly
                                    170
Ser Met Ile Ile Gly Gly Ile Asp His Ser Leu Tyr Thr Gly Ser Leu
                                185
Trp Tyr Thr Pro Ile Arg Arg Glu Trp Tyr Tyr Glu Val Ile Ile Val
                            200
Arg Val Glu Ile Asn Gly Gln Asp Leu Lys Met Asp Cys Lys Glu Tyr
                        215
                                            220
Asn Tyr Asp Lys Ser Ile Val Asp Ser Gly Thr Thr Asn Leu Arg Leu
                    230
                                        235
Pro Lys Lys Val Phe Glu Ala Ala Val Lys Ser Ile Lys Ala Ala Ser
                                    250
Ser Thr Glu Lys Phe Pro Asp Gly Phe Trp Leu Gly Glu Gln Leu Val
Cys Trp Gln Ala Gly Thr Thr Pro Trp Asn Ile Phe Pro Val Ile Ser
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280
        275
                                                285
Leu Tyr Leu Met Gly Glu Val Thr Asn Gln Ser Phe Arg Ile Thr Ile
                        295
Leu Pro Gln Gln Tyr Leu Arg Pro Val Glu Asp Val Ala Thr Ser Gln
                                        315
Asp Asp Cys Tyr Lys Phe Ala Ile Ser Gln Ser Ser Thr Gly Thr Val
                325
                                    330
Met Gly Ala Val Ile Met Glu Gly Phe Tyr Val Val Phe Asp Arg Ala
                                345
Arg Lys Arg Ile Gly Phe Ala Val Ser Ala Cys His Val His Asp Glu
                            360
                                                365
Phe Arg Thr Ala Ala Val Glu Gly Pro Phe Val Thr Leu Asp Met Glu
                        375
Asp Cys Gly Tyr Asn Ile
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Tyr Tyr Val Glu Met Thr Val Gly Ser Pro Pro Gln Thr Leu Asn Ile
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                                25
Leu Val Asp Thr Gly Ser Ser Asn Phe Ala Val Gly Ala Ala Pro His
Pro Phe Leu His Arg Tyr Tyr Gln Arg Gln Leu Ser Ser Thr Tyr Arg
Asp Leu Arg Lys Gly Val Tyr Val Pro Tyr Thr Gln Gly Lys Trp Glu
                    70
                                        75
Gly Glu Leu Gly Thr Asp Leu Val Ser Ile Pro His Gly Pro Asn Val
                                    90
Thr Val Arg Ala Asn Ile Ala Ala Ile Thr Glu Ser Asp Lys Phe Phe
           100
                                105
Ile Asn Gly Ser Asn Trp Glu Gly Ile Leu Gly Leu Ala Tyr Ala Glu
                            120
                                                125
Ile Ala Arg Pro Asp Asp Ser Leu Glu Pro Phe Phe Asp Ser Leu Val
                        135
                                            140
Lys Gln Thr His Val Pro Asn Leu Phe Ser Leu Gln Leu Cys Gly Ala
                    150
                                        155
Gly Phe Pro Leu Asn Gln Ser Glu Val Leu Ala Ser Val Gly Gly Ser
                                    170
                165
                                                        175
Met Ile Ile Gly Gly Ile Asp His Ser Leu Tyr Thr Gly Ser Leu Trp
                                185
Tyr Thr Pro Ile Arg Arg Glu Trp Tyr Tyr Glu Val Ile Ile Val Arg
                            200
                                                205
Val Glu Ile Asn Gly Gln Asp Leu Lys Met Asp Cys Lys Glu Tyr Asn
                        215
Tyr Asp Lys Ser Ile Val Asp Ser Gly Thr Thr Asn Leu Arg Leu Pro
                    230
                                        235
Lys Lys Val Phe Glu Ala Ala Val Lys Ser Ile Lys Ala Ala Ser Ser
                                    250
Thr Glu Lys Phe Pro Asp Gly Phe Trp Leu Gly Glu Gln Leu Val Cys
Trp Gln Ala Gly Thr Thr Pro Trp Asn Ile Phe Pro Val Ile Ser Leu
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```
280
        275
Tyr Leu Met Gly Glu Val Thr Asn Gln Ser Phe Arg Ile Thr Ile Leu
                        295
                                            300
Pro Gln Gln Tyr Leu Arg Pro Val Glu Asp Val Ala Thr Ser Gln Asp
                                        315
                    310
Asp Cys Tyr Lys Phe Ala Ile Ser Gln Ser Ser Thr Gly Thr Val Met
                                    330
Gly Ala Val Ile Met Glu Gly Phe Tyr Val Val Phe Asp Arg Ala Arg
                                345
Lys Arg Ile Gly Phe Ala Val Ser Ala Cys His Val His Asp Glu Phe
                            360
Arg Thr Ala Ala Val Glu Gly Pro Phe Val Thr Leu Asp Met Glu Asp
                        375
Cys Gly Tyr Asn Ile Pro
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Gly Ser Phe Val Glu Met Val Asp Asn Leu Arg Gly Lys Ser Gly Gln
                                    10
Gly Tyr Tyr Val Glu Met Thr Val Gly Ser Pro Pro Gln Thr Leu Asn
                                25
Ile Leu Val Asp Thr Gly Ser Ser Asn Phe Ala Val Gly Ala Ala Pro
His Pro Phe Leu His Arg Tyr Tyr Gln Arg Gln Leu Ser Ser Thr Tyr
                        55
Arg Asp Leu Arg Lys Gly Val Tyr Val Pro Tyr Thr Gln Gly Lys Trp
Glu Gly Glu Leu Gly Thr Asp Leu Val Ser Ile Pro His Gly Pro Asn
                                    90
Val Thr Val Arg Ala Asn Ile Ala Ala Ile Thr Glu Ser Asp Lys Phe
           100
                                105
Phe Ile Asn Gly Ser Asn Trp Glu Gly Ile Leu Gly Leu Ala Tyr Ala
                            120
Glu Ile Ala Arg Pro Asp Asp Ser Leu Glu Pro Phe Phe Asp Ser Leu
                        135
Val Lys Gln Thr His Val
145
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Gln Ser Glu Val Leu Ala Ser Val Gly Gly Ser Met Ile Ile Gly Gly
Ile Asp His Ser Leu Tyr Thr Gly Ser Leu Trp Tyr Thr Pro Ile Arg
```

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Arg Glu Trp Tyr Tyr Glu Val Ile Ile Val Arg Val Glu Ile Asn Gly
                        55
Gln Asp Leu Lys Met Asp Cys Lys Glu Tyr Asn Tyr Asp Lys Ser Ile
                    70
Val Asp Ser Gly Thr Thr Asn Leu Arg Leu Pro Lys Lys Val Phe Glu
Ala Ala Val Lys Ser Ile Lys Ala Ala Ser Ser Thr Glu Lys Phe Pro
            100
                                105
Asp Gly Phe Trp Leu Gly Glu Gln Leu Val Cys Trp Gln Ala Gly Thr
                            120
Thr Pro Trp Asn Ile Phe Pro Val Ile Ser Leu Tyr Leu Met Gly Glu
                        135
Val Thr Asn Gln Ser Phe Arg Ile Thr Ile Leu Pro Gln Gln Tyr Leu
                    150
                                        155
Arg Pro Val Glu Asp Val Ala Thr Ser Gln Asp Asp Cys Tyr Lys Phe
                                    170
Ala Ile Ser Gln Ser Ser Thr Gly Thr Val Met Gly Ala Val Ile Met
            180
                                185
Glu Gly Phe Tyr Val Val Phe Asp Arg Ala Arg Lys Arg Ile Gly Phe
                            200
                                                 205
Ala Val Ser Ala Cys His Val His Asp Glu Phe Arg Thr Ala Ala Val
                       215
Glu Gly Pro Phe Val Thr Leu Asp Met Glu Asp Cys Gly Tyr Asn Ile
                    230
                                        235
<210> 11
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Gly Ser Phe Val Glu Met Val Asp Asn Leu Arg Gly Lys Ser
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Gly Gly Ile Asp His Ser
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<211> 5
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<400> 13
Pro Tyr Thr Gln Gly
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<213> Homo sapiens

<400> 14
Ser Ser Thr Gly
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<213> Homo sapiens

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